



1600

. RAW SEQUENCE LISTING

1 <110> APPLICANT: HILLMAN, Jennifer L.

DATE: 08/28/2003 PATENT APPLICATION: US/09/768,840A TIME: 13:59:28

Input Set : A:\pf02612div_subseqlist.txt Output Set: N:\CRF4\08282003\1768840A.raw

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GOLI, Surya K.
 4 <120> TITLE OF INVENTION: CALCIUM-BINDING PROTEIN
 6 <130> FILE REFERENCE: PF-0261-2 DIV
 8 <140> CURRENT APPLICATION NUMBER: US 09/768,840A
 9 <141> CURRENT FILING DATE: 2001-01-23
11 <150> PRIOR APPLICATION NUMBER: US 09/206,499
12 <151> PRIOR FILING DATE: 1998-12-07
14 <150> PRIOR APPLICATION NUMBER: US 08/828,242
15 <151> PRIOR FILING DATE: 1997-03-31
17 <160> NUMBER OF SEQ ID NOS: 6
18 <170> SOFTWARE: PERL Program
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23 <213> ORGANISM: Homo sapiens
25 <220> FEATURE:
26 <221> NAME/KEY: misc_feature
27 <223> OTHER INFORMATION: Incyte ID No: 2509570CD1
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32 His Gly Ala Gln Gly Lys Pro Ser Pro Asp Ala Gly Pro His Gly
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34 Gln Gly Arg Val His Gln Ala Ala Pro Leu Ser Asp Ala Pro His
35
                    35
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36 Asp Asp Ala His Gly Asn Phe Gln Tyr Asp His Glu Ala Phe Leu
37
                    50
                                        55
38 Gly Arg Glu Val Ala Lys Glu Phe Asp Gln Leu Thr Pro Glu Glu
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40 Ser Gln Ala Arg Leu Gly Arg Ile Val Asp Arg Met Asp Arg Ala
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                                        85
42 Gly Asp Gly Asp Gly Trp Val Ser Leu Ala Glu Leu Arg Ala Trp
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44 Ile Ala His Thr Gln Gln Arg His Ile Arg Asp Ser Val Ser Ala
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46 Ala Trp Asp Thr Tyr Asp Thr Asp Arg Asp Gly Arg Val Gly Trp
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48 Glu Glu Leu Arg Asn Ala Thr Tyr Gly His Tyr Ala Pro Gly Glu
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50 Glu Phe His Asp Val Glu Asp Ala Glu Thr Tyr Lys Lys Met Leu
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52 Ala Arg Asp Glu Arg Arg Phe Arg Val Ala Asp Gln Asp Gly Asp
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   54 Ser Met Ala Thr Arg Glu Glu Leu Thr Ala Phe Leu His Pro Glu
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   56 Glu Phe Pro His Met Arg Asp Ile Val Ile Ala Glu Thr Leu Glu
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                      200
                                           205
                                                                210
   58 Asp Leu Asp Arg Asn Lys Asp Gly Tyr Val Gln Val Glu Glu Tyr
   59
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                                           220
                                                                225
   60 Ile Ala Asp Leu Tyr Ser Ala Glu Pro Gly Glu Glu Glu Pro Ala
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   62 Trp Val Gln Thr Glu Arg Gln Gln Phe Arg Asp Phe Arg Asp Leu
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                                           250
   64 Asn Lys Asp Gly His Leu Asp Gly Ser Glu Val Gly His Trp Val
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                                           265
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     Leu Pro Pro Ala Gln Asp Gln Pro Leu Val Glu Ala Asn His Leu
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                                                                285
   68 Leu His Glu Ser Asp Thr Asp Lys Asp Gly Arg Leu Ser Lys Ala
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   70 Glu Ile Leu Gly Asn Trp Asn Met Phe Val Gly Ser Gln Ala Thr
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  92 ggccagggga gggtgcacca ggcggccccc ctgagcgacg ctccccatga tgacgcccac 180
  93 gggaacttcc agtacgacca tgaggctttc ctgggacggg aagtggccaa ggaattcgac 240
  94 caactcacce cagaggaaag ccaggeeegt etggggegga tegtggaeeg catggaeege 300
  95 gcgggggacg gcgacggctg ggtgtcgctg gccgagcttc gcgcgtggat cgcgcacacg 360
  96 cagcagegge acataeggga eteggtgage geggeetggg acaegtaega eaeggaeege 420
  97 gacgggcgtg tgggttggga ggagctgcgc aacgccacct atggccacta cgcgcccggt 480
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98 gaagaattte atgacgtgga ggatgcagag acctacaaaa agatgctgge tegggacgag 540
99 eggegtttee gggtggeega ecaggatggg gactegatgg ecactegaga ggagetgaca 600
100 geetteetge acceegagga gtteeeteae atgegggaca tegtgattge tgaaaceetg 660
101 gaggacetgg acagaaacaa agatggetat gteeaggtgg aggagtacat egeggatetg 720
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103 egggacetee gggatetgaa caaggatggg cacetggatg ggagtgaggt gggeeaetgg 840
104 gtgetgeeee etgeeeagga ecageeeetg gtggaageea accacetget geacgagage 900
105 gacaeggaca aggatggeg getgageaaa geggaaatee tgggtaattg gaacatgttt 960

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106 gtgggcagtc aggccaccaa ctatggcgag gacctgaccc ggcaccacga tgagctgtga 1020 107 gcaccgcgca cctgccacag cctcagaggc ccgca 1055 109 <210> SEQ ID NO: 3 110 <211> LENGTH: 331 111 <212> TYPE: PRT 112 <213> ORGANISM: Homo sapiens 114 <220> FEATURE: ·115 <221> NAME/KEY: misc_feature 116 <223> OTHER INFORMATION: GenBank ID No: g1262329 118 <400> SEQUENCE: 3 119 Met Ala Arg Gly Gly Arg Gly Arg Leu Gly Leu Ala Leu Gly 120 1 121 Leu Leu Leu Ala Leu Val Leu Ala Pro Arg Val Leu Arg Ala Lys 122 123 Pro Thr Val Arg Lys Glu Arg Val Val Arg Pro Asp Ser Glu Leu 45 124 40 35 125 Gly Glu Arg Pro Pro Glu Asp Asn Gln Ser Phe Gln Tyr Asp His 50 127 Glu Ala Phe Leu Gly Lys Glu Asp Ser Lys Thr Phe Asp Gln Leu 70 128 65 129 Thr Pro Asp Glu Ser Lys Glu Arg Leu Gly Lys Ile Val Asp Arg 80 85 130 131 Ile Asp Asn Asp Gly Asp Gly Phe Val Thr Thr Glu Glu Leu Lys 100 95 133 Thr Trp Ile Lys Arg Val Gln Lys Arg Tyr Ile Phe Asp Asn Val 110 115 134 135 Ala Lys Val Trp Lys Asp Tyr Asp Arg Asp Lys Asp Asp Lys Ile 136 125 130 137 Ser Trp Glu Glu Tyr Lys Gln Ala Thr Tyr Gly Tyr Tyr Leu Gly 138 140 145 150 139 Asn Pro Ala Glu Phe His Asp Ser Ser Asp His His Thr Phe Lys 160 155 141 Lys Met Leu Pro Arg Asp Glu Arg Arg Phe Lys Ala Ala Asp Leu 142 170 175 143 Asn Gly Asp Leu Thr Ala Thr Arg Glu Glu Phe Thr Ala Phe Leu 144 185 190 145 His Pro Glu Glu Phe Glu His Met Lys Glu Ile Val Val Leu Glu 200 205 147 Thr Leu Glu Asp Ile Asp Lys Asn Gly Asp Gly Phe Val Asp Gln 215 220 148 149 Asp Glu Tyr Ile Ala Asp Met Phe Ser His Glu Glu Asn Gly Pro 230 235 150 151 Glu Pro Asp Trp Val Leu Ser Glu Arg Glu Gln Phe Asn Glu Phe 245 250 255 152 153 Arg Asp Leu Asn Lys Asp Gly Lys Leu Asp Lys Asp Glu Ile Arg 260 265 155 His Trp Ile Leu Pro Gln Asp Tyr Asp His Ala Gln Ala Glu Ala 275 280 157 Arg His Leu Val Tyr Glu Ser Asp Lys Asn Lys Asp Glu Lys Leu

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Output Set: N:\CRF4\08282003\1768840A.raw

150					290					295					300
158	mbx	T	C1	cl.,		T 011	Clu	λcn	Ψrn		Mot	Dho	Va 1	Gly	
	TIII	гуу	GIU	GIU		Leu	GIU	ASII	пр	310	Mec	FIIC	Vai	OI,	315
160		- 1 -	m1		305	61	61	3	T 0		T ***	A a n	uic	7 an	
	GIn	Ата	Thr	ASN		GIY	GIU	ASP	Leu		ьуѕ	ASII	птъ	Asp	
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179	1114	LCu	, , ,		20		5		-1-	25			_	-	30
	λνα	V = 1	17 a 1	λνα		Δen	Ser	Glu	Tien		Glu	Ara	Pro	Pro	
	ALY	Val	Val	лгу	35	пэр	DCI	·	БСС	40	Olu	*** 9			45
181) an	N an	C1 n	Cor		Cln	Titare	λan	uie		Δla	Dhe	T.On	Gly	
	ASP	ASII	GIII	ser	50	GIII	тут	азр	1113	55	AIU	riic	LCu	GIY	60
183	a 1	-		.		Dh a	3	C1 -	T 0		Dro	A cro	C1	cor	
	GIu	Asp	ser	ьуs		Pne	ASP	GIII	Leu		PIO	ASP	GIU	Ser	ду 5 75
185	_				65				_	70			3	a 1	
	GLu	Arg	Leu	GLY		TTE	vaı	Asp	Arg		Asp	ser	ASP	Gly	
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	His	Met.	Lvs				Val	Leu	Glu	Thr	Leu	Glu	Asp	Ile	Asp
203	1110	1100		0	200					205			-		210
	Lare	Δen	Glv	Asn		Phe	Val	Asp	Gln		Glu	Tvr	Ile	Ala	
205	פעם	ក្នុរា	OT 1		215			P		220		-1-			225
203	Mo+	Dha	Sor	uic		Δen	Δen	Glv	Pro		Pro	Asn	Trn	Val	
	MEL	rne	Set	nis	230	ռոհ	r S II	GIY	110	235	110	5P	P	, 41	240
207	Co-	G1	7	C1		Dha	705	λας	Dha		Δen	Lev	, Δen	Lys	
	ser	GIU	Arg	GIU		rne	ASII	wab	rne	250	vəħ	ьeu	LOII	БХЭ	255
209	0 3	.	T	3	245	3	01.	ήl.	7 ~~		m~~	т1.	T 0	Dro	
210	GŢĀ	гàг	ьeu	Asp	ьys	ASP	GIU	тте	arg	нта	ттр	тте	rea	Pro	GTII

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Input Set : A:\pf02612div_subseqlist.txt
Output Set: N:\CRF4\08282003\1768840A.raw

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214 Ser Asp Lys Asn Lys Asp Glu Met Leu Thr Lys Glu Glu Ile Leu
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                                          295
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222 <211> LENGTH: 4
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/768,840A

DATE: 08/28/2003 TIME: 13:59:29

Input Set : A:\pf02612div_subseqlist.txt
Output Set: N:\CRF4\08282003\I768840A.raw

ease Note:

e of n and/or Xaa have been detected in the Sequence Listing. Please review the quence Listing to ensure that a corresponding explanation is presented in the <220> <223> fields of each sequence which presents at least one n or Xaa.

q#:2; N Pos. 11,13,15,19